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Result
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Maximum Match 100%
Listing-first 45 summaries
EST:
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Maximum DB seq length: 2000000000
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  813.8
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                                                                                            Query
Match Length DB ID
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 29.4
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B1523145 603175911
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CNS04MA3	AA733833	AW346838	BQ339495	BQ339473	CNS03ONR	CNS05EI1	AV952826	BJ144538	BQ263135	CNS013JA	BQ284787	BI875890	BG371210	CNS01ZN3	BF515586	BG554724	BM547961	вм930281	CNS04VRK	BQ187487	AL666430	FR0052027	BB648018	AL712986	AA511690	AZ258968	CNS03Y28	BG772726	BI906774	BG293782	BG819145	B1551720	BC023215	вн349372	BF190598	B1756778	BG910325	CNS04DXR
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ALIGNMENTS

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FEATURES Source									COMMENT	JOURNAL	TITLE	AUTHORS	REFERENCE			ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION		DEFINITION	LOCUS	BI913344	
Location/Qualifiers 1887	Plate: LLAMI1613 row: m column: 05 High quality sequence stop: 782.	http://image.llnl.gov	found through the I.M.A.G.E. Consortium/LLNL at:	Clone distribution: MGC clone distribution information can be	DNA Sequencing by: Incyte Genomics. Inc.	CONA Library Arrayed by: The T.M.A.G.E. Consortium (LINI)	Tissue Procurement: Life Technologies, Inc.	Email: cgapbs-remail.nin.gov	Contact: Robert Strausberg, Ph.D.	Unpublished (1999)	National Institutes of Health, Mammalian Gene Collection (MGC)	NIH-MGC http://mgc.nci.nih.gov/.	1 (bases 1 to 887)	Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	Homo sapiens	human .	EST.	BI913344.1 GI:16177710	BI913344	mRNA sequence.	603178823F1 NIH_MGC_121 Homo sapiens cDNA clone IMAGE:5243308 5',	BI913344 887 bp mRNA linear EST 16-OCT-2001		

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BASE COUNT
ORIGIN
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                    GGTGTTATGGAGGTCAAGGTTCTGCGGACATCAGGTGCCCGGGGTACAGTCATCGTCCCC 1695
                                                                                                            GATGACCATGCAGGCATCTTCACTTTTGAATGTGATACTATTCATGTCAGTGAGAGATATT 1635
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/lab_host="DH10B"
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TITLE
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Best Local Similarity
Matches 782; Conserv
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                                                                                         ACAGGAAGCTGACTATGGAAGAAGAGGAGGCCCAAGAGGATAGCAGAGATGGGAAAGCCAG
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Plate: LLMN11605 row: f column: 03
High quality sequence stop: 790.
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Tissue Procurement: Life Technologies, Inc.
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Mammalia; Eutheria; Primates; Catarrhini;
1 (bases 1 to 791)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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/clone_lib="NIH_MGC_121"
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLN
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.lnh.gov
plate: LLAMI1605 row: f column: 03
High quality sequence stop: 778.
Location/Qualifiers
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collect Unpublished (1999)
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this is a NIH_MGC Library."
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/db_xref="taxon:9606"
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Pred. No. 1.2e-185;
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                            CTGGTACCCGGAGAACCCTTCCCTTGGGGACAAGATTGCCAGGGTCATTGTCTATTTTGT
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GCAGAACAATGAGTCCTGTTCGGGGTCATCAGACTGCAAGGAGGGTGTCATTTTGCCAAT
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Tissue Procurement: Dr. Jim Lin, University of Iowa
CDNA Library preparation: Dr. M. Bento Soares, University of Iowa
CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
CDNA Sequencing by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
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UI-M-FIO-byw-c-07-0-UI.rl NIH_BMAP_FIO Mus
IMAGE:6400062 5', mRNA sequence.
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National Institutes of Health, Mammalian
Unpublished (1999)
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Mammalia; Eutheria;
1 (bases 1 to 775)
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/dev_stage="embryo 12.5dpc"
/lab_host="bH108 (T1 phage resistant)"
/lab_host="bH108 (T1 phage resistant)
/lab
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Pred. No. 6.8e-168;
""smatches 64;
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Sciurognathi; Muridae;
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JOURNAL MEDLINE COMMENT REFERENCE AUTHORS TITLE RESULT 5 BQ189572 LOCUS SOURCE ORGANISM ACCESSION VERSION KEYWORDS DEFINITION BQ189572 665 bp mRNA lines UI-E-EJ1-aka-f-22-0-UI.rl UI-E-EJ1 Homo sapiens UI-E-EJ1-aka-f-22-0-UI 5', mRNA sequence. BQ189572 BQ189572.l GI:20365123 Program for Rat Gene Discovery and Mapping University of Iowa discovery Contact: Soares, 97044477 Normalization and subtraction: Bonaldo, M.F., Mammalia; Eukaryota; Metazoa; Homo sapiens minan (bases Res. 6 (9), Eutheria; 1 to 665) Lennon, G. and Soares, M.B 791-806 Primates; Chordata; (1996) Catarrhini; Craniata; Vertebrata; Euteleostomi; two approaches Hominidae; linear ç ar EST 30-APR-2002 CDNA clone facilitate gene

Fax: 319 335 9565
Email: msoares@blue.weeg.uiowa.edu
Email: msoares@blue.weeg.uiowa.edu
Tissue Procurement: Dr. Gregg Hageman
cDNA Library preparation: Dr. M. Bento Soares, University of
cDNA Library Arrayed by: Dr. M. Bento Soares, University of
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Researchers may obtain clones from Rese

from Research

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451 Eckstein Medical Research Building Iowa Tel: 319 335 8250 Fax: 319 335 9565

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TTGGCAACGTGACGGGCAGCAACGCCGTCAATGTCTTCCTGGGCATCGGCCTGGCCTGGCT 2503
                                                                                 TCCCAGATACGTTTGCCAGCAAAGCTGCTGCCCCCCAGGATGTATATGCAGACGCCTCCA
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Seq primer: M13 REVERSE.
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/note-"Organ: eye; vector: pT7T3-Pac (Pharmacia) with a
modified polylinker; Site_1: EcoR I; Site_2: Not I;
UI-E-EJI is a subtracted cDNA library constructed
according to Bonaldo, Lennon and Soares, Genome Research,
6:791-806, 1996. First strand cDNA synthesis was primed
with an oligo-dT primer containing a Not I site. Double
stranded cDNA was ligated to an EcoR I adaptor, digested
with Not I, and cloned directionally into pT7T3-Pac
vector. The oligonucleotide used to prime the synthesis of
first-strand cDNA contains a library tag sequence that is
located between the Not I site and the (dT)18 tail. The
sequence tags for this library are: fetal eyes, AGAATCAAGA; lens, CGATTAAGCA; eye enterior segment, AATCCAGCAT;
optic nerve, CCATTAAGTG; retina, CCGCG; Retina Foveal and
Macular, GTCC; RPE and Chorold, ACCTA. This library was
created for the program, Gene Discovery in the Visual
System, supported by National Eye Institute (NEI)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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/clone_11b="UI-E-EJ1"
/tlssue_type="fetal eyes, lens,
optic nerve, retina, Retina Fov
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Pred. No. 1.3e-163;
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Hiramoto,K., Hori,F., Ishi,Y., Ito,M., Kawai,J., Konno,H., Kouda
,M., Koya,S., Matsuyama,T., Miyazaki,A., Nomura,K., Ohno,M.,
Okazaki,Y., Okido,T., Salto,R., Sakai,C., Sakai,K., Sano,H., Sasaki
,D., Shibata,K., Shinagawa,A., Shiraki,T., Sogabe,Y., Suzuki,H.,
Tagami,M., Tagawa,A., Takahashi,F., Takeda,Y., Tanaka,T., Toya,T.,
Muramatsu,M. and Hayashizaki,Y.
MIKEN Mouse ESTS (Arakawa,T., et al. 2001)
Computer-based methods for the mouse full-length cDNA encyclopedia: real-time sequence clustering for construction of a nonredundant cDNA library. Genome Res. 11 (2), 281-289 (2001) Yamanaka, I., Kiyosawa, H., Kondo, S., Saito, T., Shinagawa, A., Aizawa, K., Fukuda, S., Hara, A., Itoh, M., Kawai, J., Shibata, K., Arakawa, T., Ishii, Y. and Hayashizaki, Y.

Mapping of 19032 mouse cDNAs on mouse chromosomes. J. Struct. Func. Genomics 2 pre, L72-L86 (2001) Please visit our web site (http://genome.gsc.riken.go.jp/) for further details.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: genome-res@gsc.riken.go.jp,
URL:http://genome.gsc.riken.go.jp/
Carninci,P., Shibata,Y., Hayatsu,N., Sugahara,Y., Shibata,K., Itoh
,M., Konno,H., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y.
Normalization and subtraction of cap-trapper-selected cDNAs to
prepare full-length cDNA libraries for rapid discovery of new
genes. Genome Res. 10 (10), 1617-1630 (2000)
wagi,K., Fujiwake,S., Inoue,K., Togawa,Y., Izawa,M., Ohara,E.,
watahiki,M., Yoneda,Y., Ishikawa,T., Ozawa,Y., Tanaka,T., Matsuura
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On Jul 9, 2000 this sequence ve
Contact: Yoshihide Hayashizaki
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The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suebiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan
Tel: 81-45-503-9222
Fax: 81-45-503-9216
                                                                                                                                                                                                                                                                                                                                                        RIKEN integrated sequence analysis (RISA) system--384-format sequencing pipeline with 384 multicapillary sequencer. Genome Res. 10 (11), 1757-1771 (2000)
Konno,H., Fukunishi,Y., Shibata,K., Itoh,M., Carninci,P., Sugahara
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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Hayashizaki,Y.
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cDNA library was prepared and sequenced in Encyclopedia Project of Genome Exploration Genomic Sciences Center and Genome Science

Mouse Genome Research Group in

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gratefully acknowledge.
121 c 203 g 1
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/clone="A930029A02"
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/dev_stage="adult"
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Roest-Crollius, H., Jaillon, O., Dasilva, C., Fizames, C.,
Bouneau, L., Billault, A., Quetier, F., Saurin, W., Bernot,
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Direct Submission
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Bernot,A., Fizames,C., Wincker,P., Brottler,P., Quetler,
Saurin,W. and Weissenbach,J.
Human gene number estimate provided by genome wide analy
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Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLN
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAML0873 row: i column: 05
High quality sequence stop: 527.
Location/Qualifiers
1. .527
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1 (bases 1 to 527)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, N
Unpublished (1999)
Contact: Robert Strausberg, Ph.I
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Tissue Procurement: David N. Louis,
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/note="Organ: brain; Vector: pCMV-SPORT6; Site_1: Not
Site_2: Sall; Cloned unidirection!... Primer: Oligo
Average insert size 2.3 kb. Constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
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/clone="IMAGE:4938316"
/clone_lib="NCI_CGAP_Brn67"
/tissue_type="anaplastic oligodendroglioma
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Pred. No. 4.2e-126;
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CCATTCCTGGAGGGACCAGTTCATGGAGGCCATCACCGTCAGTGCAGCAGGAGGATGAGGA 2126
                                                                                                                                                                                                                                                              GTTCAAGACTACGGTGGACAAACTGATCAAGAAGACAAACCTGGCCTTGGTTGTGGGGAC 2066
                                                                                                                                                                                                                                                                                                                                                                                                   AGATGTGACAGACAGGAAGCTGACTATGGAAGAAGAGGAGGCCCAAGAGGATAGCAGAGAT 274
                                     GACTGTCTTCTGGAAGGTGCTGTTTGCCTGTGTGCCCCCCACAGAGTACTGCCACGGCTG 2246
                                                                            TGAGGATGAATCCGGGGAGGAGGAGGCTG-CCTCCTGCTTTGACTACGTCATGCACTTTCT
                                                                                                 TGAGGATGAATCCGGGGAGGAGAGGCTGCCCTCCTGCTTTGACTACGTCATGCACTTCCT
                                                                                                                                                                                                                                                                                                                      GGGAAAGCCAGTATTGGGTGAACACCCCAAACTAGAAGTCATCATTGAAGAGTCCTATGA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM11487 row: b column: 13
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1 (bases 1 to 718)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-remail.nih.gov
Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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603024449F1 NIH_MGC_114 Homo
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Organ: brain; Vector: pCMV-SPORT6; Site_1: Not1; Site_2: ECORV (destroyed); RNA source anonymous pool of 6 male brains, age range 23-27 yo. Library is oligo-dr primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.5 kb, insert size range 1-3 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 019. Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5194668"
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lone IMAGE:5194668 5',
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                                                                              2321 GCTGCACCATTGGTCTCAAAGATTCAGTCACAGCTGTTGTTTTCGTGGCATTTGGCACCT
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                CTGTCCCAGATACGTTTGCCAGCAAAGCTGCTGCCCTCCAGGATGTATATGCAGACGCCT
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Tel: 402 762 4366
Fax: 402 762 4390
Email: smith@email.marc.usda.gov
Single pass sequencing. Bases called and alt_trimmed with phred
v0.980904.e. Vector identified by cross_match with the -minscore 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and -minmatch 12 options. PCR PRimers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fahrenkrug, S.C., Stone, R.T., Heat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pig.
Sus scrofa
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BF190598
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PO Box 166, Clay Center, NE 68933-0166, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Smith TPL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Design and use of two pooled tissue normalized cDNA libraries for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and Keele, J.W.
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                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                    /organism="Sus scrofa"
/db_xref="taxon:9823"
/clone_lib="MARC 2PIG"
/tissue_type="pooled"
/lab_host="BH10B"
                                                                                                                                                                                                                                                        /note-"Vector: pCMV SPORT6; Site_1: NotI; Site_2: SalI; Library made from pooled tissue from testis, ovary, endometrium, hypothalamus, pituitary, and placenta." 161 c 131 g 115 t
                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          S.C., Freking, B.A., Rohrer, G.A., Smith, T.P.L., Heaton, M.P., Grosse, W.M., Bennett, G.A., Laegr
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                                                                                                                                                                                            Score 407.8; DB 1
Pred. No. 2.7e-97;
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Laegreid,W.W.
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Sus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            A., Gebregeorgis, E., Overton, L., Russell, D., Cl
Jong, P. and Fraser, C.M.
Rat BAC End Sequences from Library CHORI-230 Eco
Unpublished (1999)
Other_GSSs: CH230-32M17.TV
Contact: Shaying Zhao
Department of Eukaryotic Genomics
The Institute for Genomic Research
9712 Medical Center Dr., Rockville, MD 20850, US
Tel: 301 838 0200
Fax: 301 838 0208
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                                                                                                                                                                                                                                                                         Email: szhaoetigr.org
Clones are derived from the rat BAC library CHORI-230
(Attp://www.chori.org/bacpac/rat230.htm). For BAC library
availability, please contact Pieter de Jong (pdejong@mail.cho.org).
Clones may be purchased from BACPAC Resources
(http://www.chori.org/bacpac/or ering_information.htm). BAC end
page: http://www.tigr.org/tdb/bac_ends/rat/bac_end_intro.html
Plate: 32 row: M column: 17
Seg pimer: SP6
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CH230-32M17.TJ CHORI-230 Segment 1
CH230-32M17, DNA sequence
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Rattus norvegicus
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                   146
           /note=""Vector: pTarBaC2.1; Site_1:
CHORI-230 Rat (BN/SsNHsd/MCW) BAC
Pieter de Jong"
123 c 124 g 106 t
                                                                                                                                                                                 /organism="Rattus norvegicus"
/strain="BN/SsNHsd/MCW"
                                                                                       /cell_type-"Brain"
                                                                                                         /sex="Female"
                                                                                                                          /clone_lib="CHORI-230 Segment
                                                                                                                                                /db_xref="taxon:10116"
/clone="CH230-32M17"
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                                                               CDNA Library Preparation: Life Technologies, Inc. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium DNA Sequencing by: Baylor College of Medicine Human Sequencing Center
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BC023215
BC023215.1 GI
Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu,
Yoon, V.S., Kowis, C.R., Lawrence,
                                                                                                                                                                                                      Submitted (04-FEB-2002) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                             Homo
                                                                                                                                 Contact: MGC help desk
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: David N.
                                           Web site: http://www.hgsc.bcm.tmc.
                                                            Center code: BCM-HGSC
                                                                                                                                                              NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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377; Conser
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D.,
cDNA Library Preparation: Michael J. Browns
Toshiyuki and Piero Carninci (RIKEN)
                                                                                     1 (bases 1 to 817)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
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/db_xref="taxon:9606"
/clone="MAGE:4932133"
/tissue_type="Brain, anaplastic ollp/19q loss"
/clone_tib="NCI_CGAP_Brn67"
/lab_host="DH10B"
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738 c 697 g 777 t
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                                                                                                                                                                                                                                                                                                         817 bp m
Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                   384
                                                                                                                                                              Craniata; Vertebrata;
Catarrhini; Hominidae
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cDNA clone IMA
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                M.D., Ph.D.
Brownstein (NHGRI),
                                                                                                          Gene
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IMAGE:5276972 5',
                                                                                                                                                                              Euteleostomi;
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                                    CCTCGCACTTCGGCTGCACCATTGGTCTCAA 2339
                                                                                                                                                                                                                                                                                                                     GA-CCAGTTCATGGAGGCCATC-ACCGTCAGTGCAGCAGGGGATG-AGGATGAGGA-TGA 2135
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                                                                                                             CTGCTTCGCCGTCTCCATCCTCATTGGCATGCTCACCGCCATCATT-GGGGACCTGG
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /tissue_type="hippocampus"
/lab_host="DH10B"
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/clone_lib="NIH_MGC_95"
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/db_xref="taxon:9606"
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93.9%;
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Pred. No. 1.3e-86;
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361
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                 CCTCTGTCCCAGATACGTTTGCCAG 2402
                                                                                 TCGGCTGCACCATTGGTCTCAAAGATTCAGTCACAGCTGTTTTTTCGTGGCATTTGGCA
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                                                                   TCGGCTGCACCATTGGTCTCAAAGATTCAGTCACAGCTGTTGTTTTCGTGGCATTTGGCA
                                                                                                                                       CCGTCTCCATCCTCATTGGCATGCTCACCGCCATCATTGGGGACCTGGCCTCGCACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tissue Procurement: David N. Louis, M.D. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 655)
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602781252F1 NCI_CGAP_Brn67 Hc
5', mRNA sequence.
BG819145
BG819145.1 GI:14166732
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Plate: LLAM10857 row:
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National Institutes of Health, Mammalian
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Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /lab_host="DH10B (T1 phage-resistant)"
/note="Organ: brain; Vector: pCMV-SPORT6; Site_1: Not1;
Site_2: Sal1; Cloned unidirectionally. Primer: Oligo dT
Average insert size 2.3 kb. Constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
a 168 c 156 g 175 t
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/clone="IMAGE:4932133"
/clone_11b="NCI_CGAP_Brn67"
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97.9%;
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Pred. No. 5.9e-85;
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CCAGTTCATGGAGGCCATCACCGTCAGTGCAGCAGGGGATGAGGATGAGGATGAATCCGG
                                                                                                                                               GGTGAACACCCCAAACTAGAAGTCATTGAA-GAGTCCTATGAGTTCAAGACTACGGT
                                                                                                                                                                                                                        GCTGACTATGGA-AGAAGAGGAGGCCAAGAGGATAGCAGAG-ATGGGAAAGCCAGTATTG
                                                                                                                                                                                                                                                                                                  GAAAACCATAAGGGTTAAAATAGTAGATGAGGAGGGAGTACGAGAGGCAAGAGAATTTCTT
                                                       GGACAAACTGATCAAGAAGACAAACCTGGCCTTGGTTGTGGGGGACCCATTCCTGGAGGGA
                                                                                                                                  GGTGAACACCCCAAACTAGAGGTCATCATTGAACGAGTCCTATGAGTTCAAGAGTACAGT
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                                                                                                                                                                                                                                                                               419;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: The Cepko Laboratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BG293782
BG293782.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          High quality sequence start: 14
High quality sequence stop: 676
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Plate: LLAM10371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (bases 1 to 677)
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                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Mus musculus"
//db_xref="taxon:10090"
//clone=="IMAGE:4502515"
//clone=_Ib="NHH_MGC_94"
//tlssue_type="retina"
//lab_host="DH10B (phage-resistant)"
//note="Organ: eye; Vector: pCMV-SPORT6; Site_1: Not1;
//note="Organ: eye; Vector: pCMV-SPORT6; Site_1: Not1;
//note="Sall; Cloned unidirectionally; oligo-dT primed.
Site_2: Sall; Cloned unidirectionally; oligo-dT primed.
Average insert size 3.3 kb. Library enriched for
full-length clones and constructed by Life Technologies.
Note: this is a NIH_MGC Library."
76 a 153 c 198 g 150 t
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